10/07/



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10 033, 195A	
Source:	OIPE	
Date Processed by STIC:	10/3/02	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.1 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- Hand Carry directly to:
 U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
 - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- 4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10/033, 195A	
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s)	
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped	
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If Intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
Invalid <213> Response	Per 1.823 of Sequence Rolles, the only valld <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 Use of <220>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
Patentin 2.0 "bug"	Please do not use "Copy to Disk" function of Patentin version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.	

AMC/MH - Biotechnology Systems Branch - 08/21/2001



OIPE

Does Not Comply Corrected Diskette Needed

RAW SEQUENCE LISTING

3 <110> APPLICANT: Fodor, Stephen P.A.

PATENT APPLICATION: US/10/033,195A

DATE: 10/03/2002

TIME: 16:19:29

Input Set : A:\2719.2002-001.txt

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Stryer, Lubert
 4
         Read, J. Leighton
 5
         Pirrung, Michael C.
 6
 8 <120> TITLE OF INVENTION: Nucleotides and Analogs Having
         Photoremovable Protecting Groups
 9
11 <130> FILE REFERENCE: 2719.2002-001
13 <140> CURRENT APPLICATION NUMBER: 10/033,195A
14 <141> CURRENT FILING DATE: 2001-12-28
16 <150> PRIOR APPLICATION NUMBER: 09/465,126
17 <151> PRIOR FILING DATE: 1999-12-17
19 <150> PRIOR APPLICATION NUMBER: 09/063,933
20 <151> PRIOR FILING DATE: 1998-04-21
22 <150> PRIOR APPLICATION NUMBER: 08/466,632
23 <151> PRIOR FILING DATE: 1995-06-06
25 <150> PRIOR APPLICATION NUMBER: 08/390,272
26 <151> PRIOR FILING DATE: 1995-02-16
28 <150> PRIOR APPLICATION NUMBER: 07/624,120
29 <151> PRIOR FILING DATE: 1990-12-06
31 <150> PRIOR APPLICATION NUMBER: 07/492,462
32 <151> PRIOR FILING DATE: 1990-03-07
34 <150> PRIOR APPLICATION NUMBER: 07/362,901
35 <151> PRIOR FILING DATE: 1989-06-07
37 <150> PRIOR APPLICATION NUMBER: 08/456,887
38 <151> PRIOR FILING DATE: 1995-06-01
                                                          The type of errors shown exist throughout
                                                         the Sequence Listing. Please check subsequent
40 <150> PRIOR APPLICATION NUMBER: 07/954,646
41 <151> PRIOR FILING DATE: 1992-09-30
                                                  must explain genetic source

See ever summany sheet item 11
                                                         sequences for similar errors.
43 <150> PRIOR APPLICATION NUMBER: 07/850,356
44 <151> PRIOR FILING DATE: 1992-03-12
46 <160> NUMBER OF SEQ ID NOS: 20
48 <170> SOFTWARE: FastSEQ for Windows Version 4.0
50 <210> SEO ID NO: 1
51 <211> LENGTH: 5
52 <212> TYPE: PRT
53 <213> ORGANISM: Artificial Sequence
55 <220> FEATURE:
56 <223> OTHER INFORMATION ( Peptide sequence
60 <400> SEQUENCE: 1
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62 1
65 <210> SEQ ID NO: 2
66 <211> LENGTH: 5
67 <212> TYPE: PRT
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PATENT APPLICATION: US/10/033,195A

DATE: 10/03/2002 TIME: 16:19:29

Input Set : A:\2719.2002-001.txt

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70 <220> FEATURE:
71 <223> OTHER INFORMATION: Peptide sequence
73 <400> SEQUENCE: 2
74 Pro Gly Gly Phe Leu
75 1
78 <210> SEQ ID NO: 3
79 <211> LENGTH: 6
80 <212> TYPE: PRT
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION:
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86 <400> SEQUENCE: 3
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92 <211> LENGTH: 5
93 <212> TYPE: PRT
94 <213> ORGANISM: Artificial Sequence
96 <220> FEATURE:
97 <223> OTHER INFORMATION ( Peptide sequence
99 <400> SEQUENCE: 4
100 Tyr Gly Ala Phe Ser
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104 <210> SEQ ID NO: 5
105 <211> LENGTH: 5
106 <212> TYPE: PRT
107 <213> ORGANISM: Artificial Sequence
109 <220> FEATURE:
110 <223> OTHER INFORMATION: Peptide sequence
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117 <210> SEQ ID NO: 6
118 <211> LENGTH: 6
119 <212> TYPE: PRT
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127 1
130 <210> SEQ ID NO: 7
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132 <212> TYPE: PRT
133 <213> ORGANISM: Artificial Sequence
135 <220> FEATURE:
136 <223> OTHER INFORMATION: Peptide sequence
138 <400> SEQUENCE: 7
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DATE: 10/03/2002 TIME: 16:19:29 PATENT APPLICATION: US/10/033,195A

Input Set : A:\2719.2002-001.txt

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149 <223> OTHER INFORMATION: Peptide sequence
151 <400> SEQUENCE: 8
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156 <210> SEQ ID NO: 9
157 <211> LENGTH: 5
158 <212> TYPE: PRT
159 <213> ORGANISM: Artificial Sequence
161 <220> FEATURE:
162 <223> OTHER INFORMATION: Peptide sequence
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165 Tyr Gly Gly Phe Ser
166
169 <210> SEQ ID NO: 10
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183 <210> SEQ ID NO: 11
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204 <223> OTHER INFORMATION: Peptide sequence
206 <400> SEQUENCE: 12
207 Tyr Gly Ala Phe Phe
208 1
211 <210> SEQ ID NO: 13
212 <211> LENGTH: 5
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DATE: 10/03/2002 TIME: 16:19:29 PATENT APPLICATION: US/10/033,195A

Input Set : A:\2719.2002-001.txt

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214 <213> ORGANISM: Artificial Sequence
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217 <223> OTHER INFORMATION: Peptide sequence
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224 <210> SEQ ID NO: 14
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226 <212> TYPE: PRT
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229 <220> FEATURE:
230 <223> OTHER INFORMATION: Peptide sequence
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233 Tyr Gly Ala Phe Ser Phe
234 1
237 <210> SEQ ID NO: 15
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239 <212> TYPE: PRT
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250 <210> SEQ ID NO: 16
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265 <212> TYPE: PRT
266 <213> ORGANISM: Artificial Sequence
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277 <211> LENGTH: 5
278 <212> TYPE: PRT
279 <213> ORGANISM: Artificial Sequence
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DATE: 10/03/2002 TIME: 16:19:29

Input Set : A:\2719.2002-001.txt

PATENT APPLICATION: US/10/033,195A

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295 <223> OTHER INFORMATION: Peptide sequence
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303 <211> LENGTH: 4
304 <212> TYPE: PRT
305 <213> ORGANISM: Artificial Sequence
307 <220> FEATURE:
308 <223> OTHER INFORMATION: Peptide sequence
311 <400> SEQUENCE: 20
312 Gly Gly Phe Leu
313 1
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VERIFICATION SUMMARY

DATE: 10/03/2002 TIME: 16:19:30

PATENT APPLICATION: US/10/033,195A

Input Set : A:\2719.2002-001.txt